

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/532,449A
Source: 1FW/6
Date Processed by STIC: 10/5/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/efc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER:

10/537,449A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped
 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000

- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence. (see item 11 below)

- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules

- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

- 13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



IFW16

RAW SEQUENCE LISTING

DATE: 10/05/2006

PATENT APPLICATION: US/10/537,449A

TIME: 11:38:44

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\10052006\J537449A.raw

3 <110> APPLICANT: Technische Universitat Dresden
 5 <120> TITLE OF INVENTION: Polynucleotides Targeted Against Htert and Use Thereof
 7 <130> FILE REFERENCE: 101215-189-2
 9 <140> CURRENT APPLICATION NUMBER: 10/537,449A
 C--> 10 <141> CURRENT FILING DATE: 2006-01-09
 12 <160> NUMBER OF SEQ ID NOS: 18
 14 <170> SOFTWARE: PatentIn Ver. 2.1
 16 <210> SEQ ID NO: 1
 17 <211> LENGTH: 75
 18 <212> TYPE: DNA
 19 <213> ORGANISM: Homo sapiens
 21 <220> FEATURE:
 22 <221> NAME/KEY: mRNA
 23 <222> LOCATION: (1)..(75)
 24 <223> OTHER INFORMATION: subunit 2176-2250 of hTERT
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 27 <308> DATABASE ACCESSION NO: AF015950
 W--> 28 <309> DATABASE ENTRY DATE: AUG 1997 *AUG-1997 (global env)*
 30 <400> SEQUENCE: 1
 31 ctttgtcaag gtggatgtga cgggcgcgta cgacaccatc cccaggaca ggctcacgga 60
 32 ggtcatcgcc agcat 75
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 36 <211> LENGTH: 98
 37 <212> TYPE: DNA
 38 <213> ORGANISM: Homo sapiens
 40 <220> FEATURE:
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 42 <222> LOCATION: (1)..(98)
 43 <223> OTHER INFORMATION: subunit 2296-2393 of hTERT
 W--> 45 <307> DATE:
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 50 ccagaaggcc gcccatgggc acgtccgcaa ggccttcaag agccacgtct ctaccttgac 60
 51 agacctccag ccgtacatgc gacagttcgt ggctcacc 98
 54 <210> SEQ ID NO: 3
 55 <211> LENGTH: 23
 56 <212> TYPE: DNA
 57 <213> ORGANISM: Homo sapiens
 59 <220> FEATURE:
 60 <221> NAME/KEY: mRNA
 61 <222> LOCATION: (1)..(23)
 62 <223> OTHER INFORMATION: subunit 2183-2205 of hTERT

Does Not Comply
 Corrected Diskette Needed

insert this mandatory human identifier above publication information. <307> is a "header" only. It never has a response.

RAW SEQUENCE LISTING

DATE: 10/05/2006

PATENT APPLICATION: US/10/537,449A

TIME: 11:38:44

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\10052006\J537449A.raw

insert
↓
<3007

W--> 64 ~~<307> DATE:~~
65 <308> DATABASE ACCESSION NO: AF015950
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69 aaggtggatg tgacgggcgc gta 23
72 <210> SEQ ID NO: 4
73 <211> LENGTH: 20
74 <212> TYPE: DNA
75 <213> ORGANISM: Homo sapiens
77 <220> FEATURE:
78 <221> NAME/KEY: mRNA
79 <222> LOCATION: (1)..(20)
80 <223> OTHER INFORMATION: subunit 2206-2225 of hTERT *<3007*

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83 <308> DATABASE ACCESSION NO: AF015950
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86 <400> SEQUENCE: 4
87 cgacaccatc ccccaggaca 20
90 <210> SEQ ID NO: 5
91 <211> LENGTH: 20
92 <212> TYPE: DNA
93 <213> ORGANISM: Homo sapiens
95 <220> FEATURE:
96 <221> NAME/KEY: mRNA
97 <222> LOCATION: (1)..(20)
98 <223> OTHER INFORMATION: subunit 2315-2334 of hTERT *<3007*

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W--> 102 <309> DATABASE ENTRY DATE: AUG 1997
104 <400> SEQUENCE: 5
105 cacgtccgca aggccttcaa 20
108 <210> SEQ ID NO: 6
109 <211> LENGTH: 20
110 <212> TYPE: DNA
111 <213> ORGANISM: Homo sapiens
113 <220> FEATURE:
114 <221> NAME/KEY: mRNA
115 <222> LOCATION: (1)..(20)
116 <223> OTHER INFORMATION: subunit 2317-2336 of hTERT *<3007*

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122 <400> SEQUENCE: 6
123 cgtccgcaag gccttcaaga 20
126 <210> SEQ ID NO: 7
127 <211> LENGTH: 23
128 <212> TYPE: DNA
129 <213> ORGANISM: Homo sapiens
131 <220> FEATURE:

RAW SEQUENCE LISTING

DATE: 10/05/2006

PATENT APPLICATION: US/10/537,449A

TIME: 11:38:44

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\10052006\J537449A.raw

132 <221> NAME/KEY: mRNA
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141 aaggccttca agagccacgt etc. 23
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147 <213> ORGANISM: Homo sapiens
149 <220> FEATURE:
150 <221> NAME/KEY: mRNA
151 <222> LOCATION: (1)..(20)
152 <223> OTHER INFORMATION: subunit 2331-2350 hTERT (3007)
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158 <400> SEQUENCE: 8 20
159 tcaagagcca cgtctctacc
162 <210> SEQ ID NO: 9
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164 <212> TYPE: DNA
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170 <223> OTHER INFORMATION: subunit 2333-2352 of hTERT (3007)
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176 <400> SEQUENCE: 9 20
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182 <212> TYPE: DNA
183 <213> ORGANISM: Artificial Sequence
185 <220> FEATURE:
186 <223> OTHER INFORMATION: hTERT-AS AStel 2206-2225 (3007)
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189 <308> DATABASE ACCESSION NO: AF015950
W--> 190 <309> DATABASE ENTRY DATE: AUG 1997
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196 <210> SEQ ID NO: 11
197 <211> LENGTH: 20
198 <212> TYPE: DNA
199 <213> ORGANISM: Artificial Sequence

RAW SEQUENCE LISTING

DATE: 10/05/2006

PATENT APPLICATION: US/10/537,449A

TIME: 11:38:44

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\10052006\J537449A.raw

201 <220> FEATURE:
 202 <223> OTHER INFORMATION: hTERT-AS AStel 2315-2334 (3007)
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 205 <308> DATABASE ACCESSION NO: AF015950
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 208 <400> SEQUENCE: 11
 209 ttgaaggcct tgcggacgtg 20
 212 <210> SEQ ID NO: 12
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 214 <212> TYPE: DNA
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 217 <220> FEATURE:
 218 <223> OTHER INFORMATION: hTERT-AS AStel 2317-2336 (3007)
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 221 <308> DATABASE ACCESSION NO: AF015950
 W--> 222 <309> DATABASE ENTRY DATE: AUG 1997
 224 <400> SEQUENCE: 12
 225 tcttgaaggc cttgcggacg 20
 228 <210> SEQ ID NO: 13
 229 <211> LENGTH: 20
 230 <212> TYPE: DNA
 231 <213> ORGANISM: Artificial Sequence
 233 <220> FEATURE:
 234 <223> OTHER INFORMATION: hTERT-AS AStel 2331-2350 (3007)
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 237 <308> DATABASE ACCESSION NO: AF015950
 W--> 238 <309> DATABASE ENTRY DATE: AUG 1997
 240 <400> SEQUENCE: 13
 241 ggtagagacg tggctcttga 20
 244 <210> SEQ ID NO: 14
 245 <211> LENGTH: 20
 246 <212> TYPE: DNA
 247 <213> ORGANISM: Artificial Sequence
 249 <220> FEATURE:
 250 <223> OTHER INFORMATION: hTERT-AS AStel 2333-2352 (3007)
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 253 <308> DATABASE ACCESSION NO: AF015950
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 260 <210> SEQ ID NO: 15
 261 <211> LENGTH: 20
 262 <212> TYPE: DNA
 263 <213> ORGANISM: Artificial Sequence
 265 <220> FEATURE:
 266 <223> OTHER INFORMATION: NS-K2 *what is its source? (see item 11 on*
 268 <300> PUBLICATION INFORMATION: *Erra Summary*
 269 <308> DATABASE ACCESSION NO: AF015950
 W--> 270 <309> DATABASE ENTRY DATE: AUG 1997 *Sheet)*

RAW SEQUENCE LISTING

DATE: 10/05/2006

PATENT APPLICATION: US/10/537,449A

TIME: 11:38:44

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\10052006\J537449A.raw

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272 <400> SEQUENCE: 15
273 cagtctcagt actgaagctg 20
276 <210> SEQ ID NO: 16
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278 <212> TYPE: DNA
279 <213> ORGANISM: Artificial Sequence
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284 <300> PUBLICATION INFORMATION:
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289 cagcttcagt actgagactg 20
292 <210> SEQ ID NO: 17
293 <211> LENGTH: 501
294 <212> TYPE: DNA
295 <213> ORGANISM: Homo sapiens
297 <220> FEATURE:
298 <221> NAME/KEY: mRNA
299 <222> LOCATION: (1)..(501)
300 <223> OTHER INFORMATION: subunit 2000-2500 of hTERT
302 <300> PUBLICATION INFORMATION:
303 <308> DATABASE ACCESSION NO: AF015950
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306 <400> SEQUENCE: 17
307 aagagggccg agcgtctcac ctcgaggggtg aaggcactgt tcagcgtgct caactacgag 60
308 cgggcgcggc gcccggcct cctgggcgcc tctgtgctgg gcctggacga tatccacagg 120
309 gcctggcgca ccttcgtgct gcgtgtgcgg gcccaggacc cgccgcctga gctgtacttt 180
310 gtcaagggtg atgtgacggg cgcgtacgac accatcccc aggacaggct cacggaggctc 240
311 atcgccagca tcatcaaacc ccagaacacg tactgcgtgc gtcggtatgc cgtggtccag 300
312 aaggccgccc atgggcacgt ccgcaaggcc ttcaagagcc acgtctctac cttgacagac 360
313 ctccagccgt acatgcgaca gtctgtggct cacctgcagg agaccagccc gctgagggat 420
314 gccgtcgtca tcgagcagag ctctccctg aatgaggcca gcagtggcct cttcgacgtc 480
315 ttctacgct tcatgtgcca c 501
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324 <221> NAME/KEY: mRNA
325 <222> LOCATION: (1)..(4015)
326 <223> OTHER INFORMATION: hTERT
328 <300> PUBLICATION INFORMATION:
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332 <400> SEQUENCE: 18
333 gcagcgtgc gtctgtgc gcacgtggga agccctggcc ccggccaccc ccgcgatgcc 60
334 gcgcgtccc cgctgccgag ccgtgcgtc cctgctgcgc agccactacc gcgaggtgct 120
335 gccgctggcc acgttcgtgc ggcgctggg gcccagggc tggcggtgg tgcagcgcg 180

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VERIFICATION SUMMARY

DATE: 10/05/2006

PATENT APPLICATION: US/10/537,449A

TIME: 11:38:45

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\10052006\J537449A.raw

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:26 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:1
L:28 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:1
L:45 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:2
L:47 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:2
L:64 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:3
L:66 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:3
L:82 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:4
L:84 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:4
L:100 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:5
L:102 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:5
L:118 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:6
L:120 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:6
L:136 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:7
L:138 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:7
L:154 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:8
L:156 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:8
L:172 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:9
L:174 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:9
L:188 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:10
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L:204 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:11
L:206 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:11
L:220 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:12
L:222 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:12
L:236 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:13
L:238 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:13
L:252 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:14
L:254 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:14
L:270 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:15
L:286 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:16
L:304 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:17
L:330 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:18